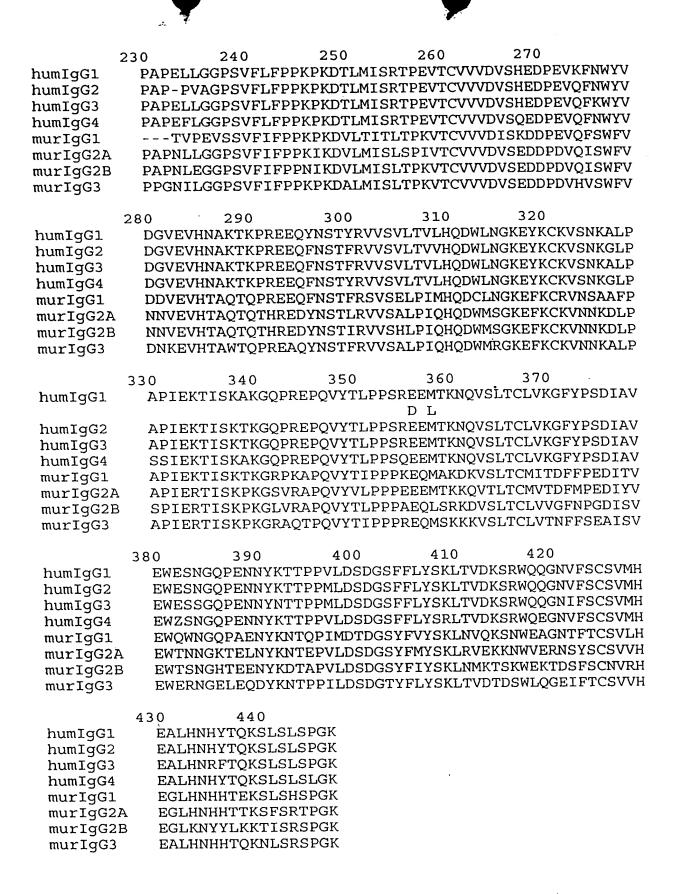
Fig.1





## Percent Identity Among Fc Sequences

	•	1	2	3	4	5	6	7	8
1.	humIgG1	-	94	94	94	64	66	63	68
2.	humIgG2		-	93	92	65	63	60	67
3.	humIgG3			-	91	64	64	61	67
4.	humIgG4				-	62	64	61	64
5.	murIgGl					-	65	61	67
6.	murIgG2A						-	77	70
7.	murIgG2B							-	68
Я	murTaG3								-

Fig. 2B

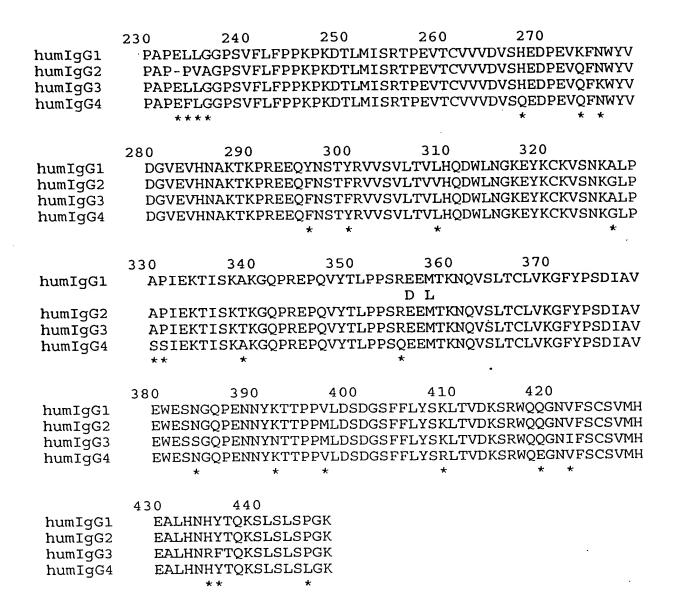
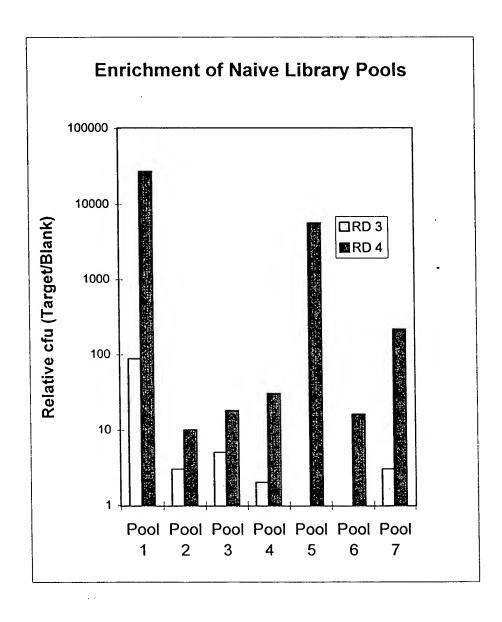


Fig. 3



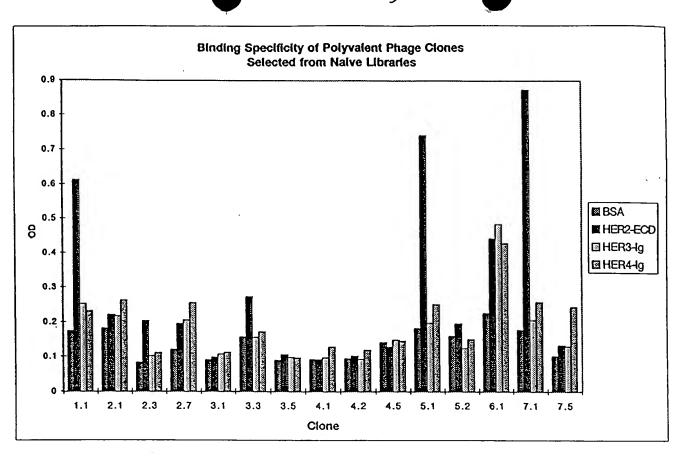
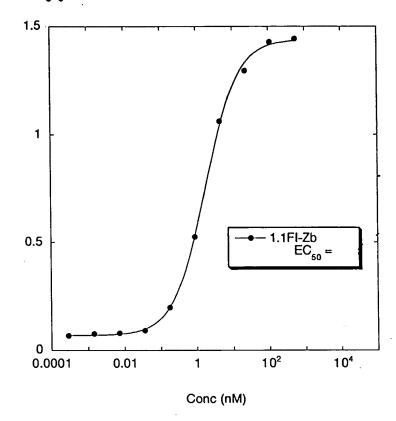


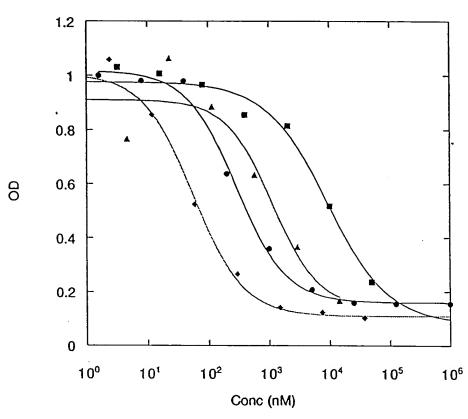
Figure 6.

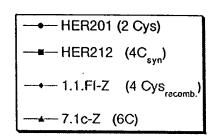
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## $EC_{5\ 0}$ for 1.1.Fl-Zb Binding to Immobilized HER2

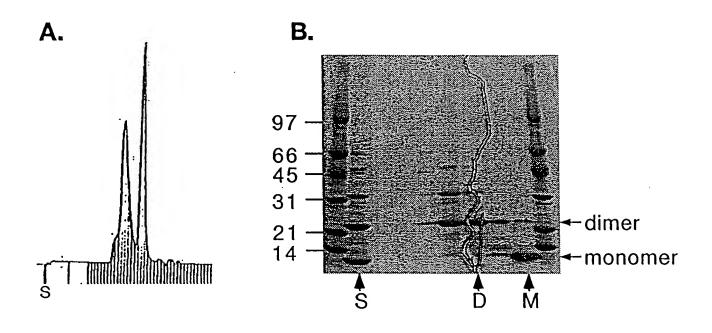


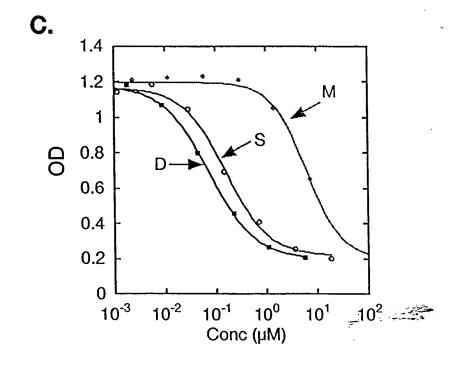






1.1.Fl-Z
QVYESWGCIGPGCACLQACL-Z

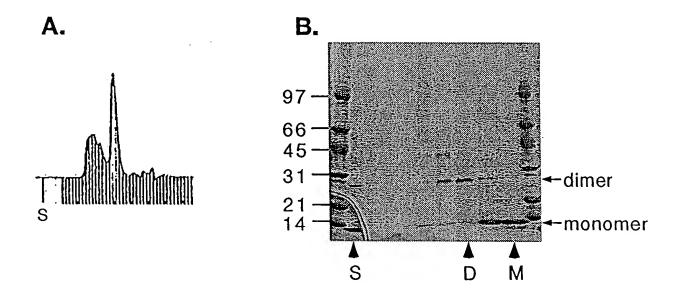




ot <del>i</del> ...

 $(1.1.FI)_2$ -Z

## QVYESWGCIGPGCACLQACLGGGSGGQVYESWGCIGPGCACLQACL-Z



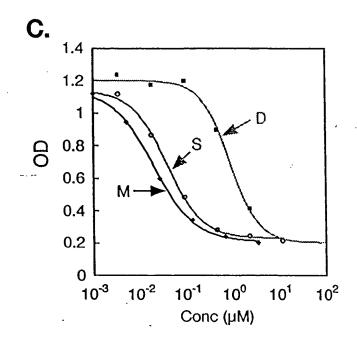
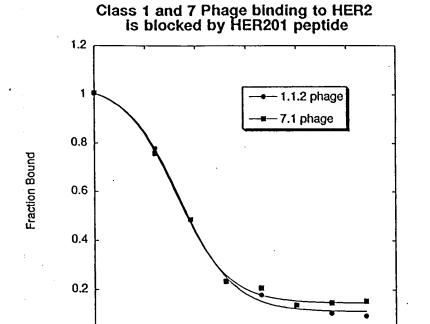


Figure (0



0

1

10¹

 $10^2$ 



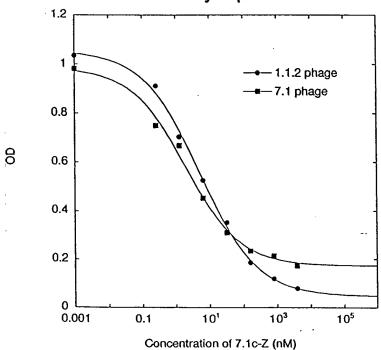
 $10^{3}$ 

Concentration of HER201 (nM)

10⁴

10<sup>5</sup>

10<sup>6</sup> -

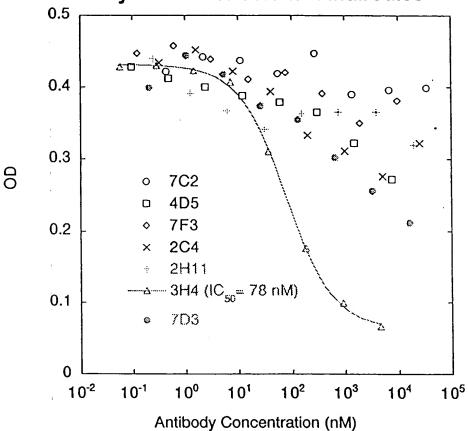


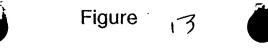
\*\*\*\*\* \

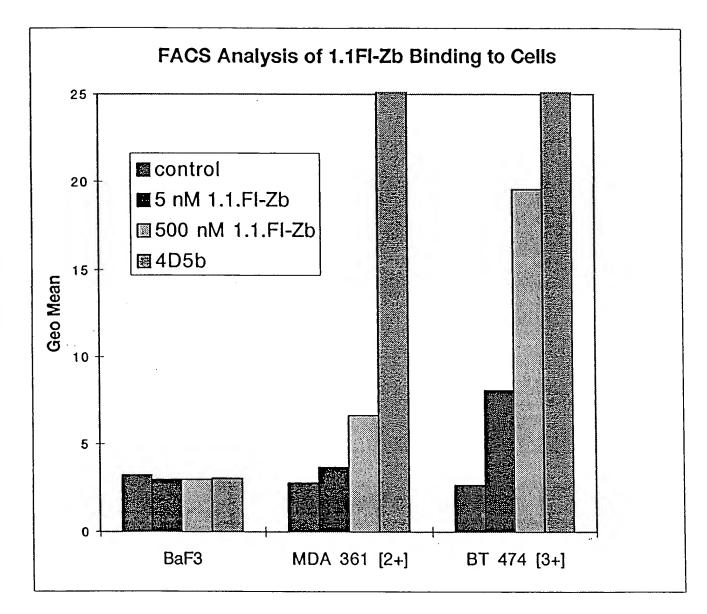
Class 7

CGGEWWHCCGPGCGWVVDAC 7.1c Y S F E G W G C I G P G C A Y L F E G H 1.1.2 Y E W E G W G C I G P G C P A L G F G Y 1.1.FC EAWGCIGPGCEMLGAWC 1.1.FA Class 1 ERWGCIGPGCECLMSCV 1.1.FB C I D E T W G C I G P G C E E L R C K R 1.1.FH FWGCIGPGCAQL 1.1.CF

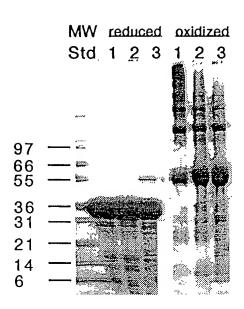








.\*:•



1 = Control Fc (no peptide fusion) 2 = HER2-Fc fusion (1.1 FI-Fc) F16UZE Sequence

15

clone

## Pool 1

1.1

49-9-4-N-0210

Y E V E A W D G M G P G O A N L F E A H

Library 1.1
1.1.8
1.1.9
1.1.10
1.1.1
1.1.2
1.1.6
1.1.11
1.1.3
1.1.4
1.1.5

 Y E F E G W D C M G P G C A S V F G A H

 Y E G E S W G C I G P G C A F L E G A H

 Y E V F G W G C M G P G C A F L L E A H

 Y E V F G W G C M G P G C A F L L E A H

 Y D F E G W G C I G P G C A Y L F G A H

 Y D F E G W G C I G P G C A Y L F G A H

 Y D F E G W G C I G P G C A Y L F G A L

 Y D F E G W G C I G P G C A Y L F E A Y

 F E F E G W G C I G P G C A Y L F E A Y

 F E F E G W G C I G P G C A D L F E A F

YEFEGWGO I GROCAYLFEAH

Pool 7

1.1.7

Consensus

7.1 SSWVLVQGGGEWWHGGGJGGGLVVNA

Library 7.1		
7.1.1	GAWVSFEGGGEVWAAGGGEGWVWKAX	
7.1.9	CAWYLVOCGGEWWHCCGGPGGEFVVDA	2
7.1.6	GAWXAVYCGGELWHGCGGPGCGFVYDS	8
7.1.5	EAWVRVWCFGEWWDCCGLGCGWVV <u>N</u> V	3
7.1.3	GAWVRVLCGGEWWHCGGGLG@GWVEVEA.	3
7.1.4	CSWVSVLCGGEWWQCCGPGCGLVVNA	
7.1.2	6 S W V S L 6 6 G G E W W H 6 6 G G G 6 W V V N V	
7.1.7	CSWVLLHCGGEWWHCCGGGGGGWVGEA	9
7.1.8	CISWVLLECGGELWEGGGEGGWVADA	Ò
7.1.11	CSWVVFECLGESWHCCGGGWVVHA	0
Consensus	CAWVSVECGGEWWHOCGPGCGWVVDA	Ō



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Figure 12

